

AMENDMENTS TO THE SPECIFICATION:

Please amend the paragraph on page 12, line 24 – page 13, line 2 as follows:

The “Carbohydrate-Binding Module of Family 20” or a CBM-20 module is in the context of this invention defined as a sequence of approximately 100 amino acids having at least 45% identity to the Carbohydrate-Binding Module (CBM) of the polypeptide disclosed in figure 1 by Joergensen et al (1997) in *Biotechnol. Lett.* 19:1027-1031. The CBM comprises the last 102 amino acids of the polypeptide, i.e., the subsequence from amino acid 582 to amino acid 683. The numbering of Glycoside Hydrolase Families applied in this disclosure follows the concept of Coutinho, P.M. & Henrissat, B. (1999) *CAZy - Carbohydrate-Active Enzymes server* at URL: ~~afmb.cnrs-mrs.fr/~cazy/CAZY/index.html~~ on the internet or alternatively Coutinho, P.M. & Henrissat, B. 1999; The modular structure of cellulases and other carbohydrate-active enzymes: an integrated database approach. In *"Genetics, Biochemistry and Ecology of Cellulose Degradation"*, K. Ohmiya, K. Hayashi, K. Sakka, Y. Kobayashi, S. Karita and T. Kimura eds., Uni Publishers Co., Tokyo, pp. 15-23, and Bourne, Y. & Henrissat, B. 2001; Glycoside hydrolases and glycosyltransferases: families and functional modules, *Current Opinion in Structural Biology* 11:593-600.

Please amend the paragraph on page 14, lines 28-29 as follows:

Further suitable CBMs of Carbohydrate-Binding Module Family 20 may be found on the internet at URL: ~~afmb.cnrs-mrs.fr/~cazy/CAZY/index.html~~).

Please amend the paragraph on page 23, lines 34-35 as follows:

A two dose dose-response was conducted with each enzyme. Dosages used were ~~0.3 and 0.6 nmol/g DS~~ 0.3 and 0.6 nmol/g DS. Six replicates of each treatment were run.